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&lt;170&gt; PatentIn Ver. 2.1

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Trp	Ser	Ser	Cys	Ser	Arg	Asp	Tyr	Ile	Thr	Ser	Phe	Leu	Asp	Ser	Gly
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Leu	Gly	Leu	Cys	Leu	Asn	Asn	Arg	Pro	Pro	Arg	Gln	Asp	Phe	Val	Tyr
	450					455					460				
Pro	Thr	Val	Ala	Pro	Gly	Gln	Ala	Tyr	Asp	Ala	Asp	Glu	Gln	Cys	Arg
465					470					475					480
Phe	Gln	His	Gly	Val	Lys	Ser	Arg	Gln	Cys	Lys	Tyr	Gly	Glu	Val	Cys
				485					490					495	
Ser	Glu	Leu	Trp	Cys	Leu	Ser	Lys	Ser	Asn	Arg	Cys	Ile	Thr	Asn	Ser
		500						505					510		
Ile	Pro	Ala	Ala	Glu	Gly	Thr	Leu	Cys	Gln	Thr	His	Thr	Ile	Asp	Lys
		515					520					525			
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	530					535					540				
Gly	Val	Asp	Gly	Ala	Trp	Gly	Pro	Trp	Thr	Pro	Trp	Gly	Asp	Cys	Ser
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Arg	Ser	Cys	Gly	Gly	Gly	Val	Ser	Ser	Ser	Ser	Arg	His	Cys	Asp	Ser
				565					570					575	
Pro	Arg	Pro	Thr	Ile	Gly	Gly	Lys	Tyr	Cys	Leu	Gly	Glu	Arg	Arg	Arg
			580					585					590		
His	Arg	Ser	Cys	Asn	Thr	Asn	Asp	Cys	Pro	Pro	Gly	Ser	Gln	Asp	Phe
		595					600					605			



Arg 610	Glu	Met	Gln	Cys	Ser	Glu 615	Phe	Asp	Ser	Val	Pro 620	Phe	Arg	Gly	Lys
Phe 625	Tyr	Thr	Trp	Lys	Thr 630	Tyr	Arg	Gly	Gly	Gly 635	Val	Lys	Ala	Cys	Ser 640
Leu	Thr	Cys	Leu	Ala 645	Glu	Gly	Phe	Asn	Phe 650	Tyr	Thr	Glu	Arg	Ala 655	Ala
Ala	Val	Val	Asp 660	Gly	Thr	Pro	Cys	Arg 665	Pro	Asp	Thr	Val	Asp 670	Ile	Cys
Val	Ser	Gly 675	Glu	Cys	Lys	His	Val 680	Gly	Cys	Asp	Arg	Val 685	Leu	Gly	Ser
Asp 690	Leu	Arg	Glu	Asp	Lys	Cys 695	Arg	Val	Cys	Gly	Gly 700	Asp	Gly	Ser	Ala
Cys 705	Glu	Thr	Ile	Glu	Gly 710	Val	Phe	Ser	Pro	Ala 715	Leu	Pro	Gly	Thr	Gly 720
Tyr	Glu	Asp	Val	Val 725	Trp	Ile	Pro	Lys	Gly 730	Ser	Val	His	Ile	Phe 735	Ile
Gln	Asp	Leu	Asn 740	Leu	Ser	Leu	Ser	His 745	Leu	Ala	Leu	Lys	Gly 750	Asp	Gln
Glu	Ser	Leu 755	Leu	Leu	Glu	Gly	Leu 760	Pro	Gly	Thr	Pro	Gln 765	Pro	His	Arg
Leu 770	Pro	Leu	Ala	Gly	Thr	Thr 775	Phe	His	Leu	Arg	Gln 780	Gly	Pro	Asp	Gln
Ala 785	Gln	Ser	Leu	Glu	Ala 790	Leu	Gly	Pro	Ile	Asn 795	Ala	Ser	Leu	Ile	Ile 800
Met	Val	Leu	Ala	Gln 805	Ala	Glu	Leu	Pro	Ala 810	Leu	His	Tyr	Arg	Phe 815	Asn
Ala	Pro	Ile	Ala 820	Arg	Asp	Ala	Leu	Pro 825	Pro	Tyr	Ser	Trp	His 830	Tyr	Ala
Pro	Trp	Thr 835	Lys	Cys	Ser	Ala	Gln 840	Cys	Ala	Gly	Gly	Ser 845	Gln	Val	Gln
Val 850	Val	Glu	Cys	Arg	Asn	Gln 855	Leu	Asp	Ser	Ser	Ala 860	Val	Ala	Pro	His
Tyr 865	Cys	Ser	Gly	His	Ser 870	Lys	Leu	Pro	Lys	Arg 875	Gln	Arg	Ala	Cys	Asn 880
Thr	Glu	Pro	Cys	Pro 885	Pro	Asp	Trp	Val	Val 890	Gly	Asn	Trp	Ser	Arg 895	Cys
Ser	Arg	Ser	Cys 900	Asp	Ala	Gly	Val	Arg 905	Ser	Arg	Ser	Val 910	Val	Cys	Gln
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Protein Data Bank

Pro Gln Pro Arg Pro Pro Val Leu Glu Ala Cys Gln Gly Pro Met Cys  
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Pro Pro Glu Trp Ala Thr Leu Asp Trp Ser Glu Cys Thr Pro Ser Cys  
945 950 955 960

Gly Pro Gly Leu Arg His Arg Val Val Leu Cys Lys Ser Ala Asp Gln  
965 970 975

Arg Ser Thr Leu Pro Pro Gly His Cys Leu Pro Ala Ala Lys Pro Pro  
980 985 990

Ser Thr Met Arg Cys Asn Leu Arg Arg Cys Pro Pro Ala Arg Trp Val  
995 1000 1005

Thr Ser Glu Trp Gly Glu Cys Ser Thr Gln Cys Gly Leu Gly Gln Gln  
1010 1015 1020

Gln Arg Thr Val Arg Cys Thr Ser His Thr Gly Gln Pro Ser Arg Glu  
1025 1030 1035 1040

Cys Thr Glu Ala Leu Arg Pro Ser Thr Met Gln Gln Cys Glu Ala Lys  
1045 1050 1055

Cys Asp Ser Val Val Pro Pro Gly Asp Gly Pro Glu Glu Cys Lys Asp  
1060 1065 1070

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His Val Val Tyr Lys Arg Ser Ser Leu Arg His Pro His Leu Asp Thr  
180 185 190

Ala Cys Gly Val Arg Asp Glu Lys Pro Trp Lys Gly Arg Pro Trp Trp  
195 200 205

Leu Arg Thr Leu Lys Pro Pro Pro Ala Arg Pro Leu Gly Asn Glu Thr  
210 215 220

Glu Arg Gly Gln Pro Gly Leu Lys Arg Ser Val Ser Arg Glu Arg Tyr  
225 230 235 240

Val Glu Thr Leu Val Val Ala Asp Lys Met Met Val Ala Tyr His Gly  
245 250 255

Arg Arg Asp Val Glu Gln Tyr Val Leu Ala Ile Met Asn Ile Val Ala  
260 265 270

Lys Leu Phe Gln Asp Ser Ser Leu Gly Ser Thr Val Asn Ile Leu Val  
275 280 285

Thr Arg Leu Ile Leu Leu Thr Glu Asp Gln Pro Thr Leu Glu Ile Thr  
290 295 300

His His Ala Gly Lys Ser Leu Asp Ser Phe Cys Lys Trp Gln Lys Ser  
305 310 315 320

Ile Val Asn His Ser Gly His Gly Asn Ala Ile Pro Glu Asn Gly Val  
325 330 335

Ala Asn His Asp Thr Ala Val Leu Ile Thr Arg Tyr Asp Ile Cys Ile  
340 345 350

Tyr Lys Asn Lys Pro Cys Gly Thr Leu Gly Leu Ala Pro Val Gly Gly  
355 360 365

Met Cys Glu Arg Glu Arg Ser Cys Ser Val Asn Glu Asp Ile Gly Leu  
370 375 380

Ala Thr Ala Phe Thr Ile Ala His Glu Ile Gly His Thr Phe Gly Met  
385 390 395 400

Asn His Asp Gly Val Gly Asn Ser Cys Gly Ala Arg Gly Gln Asp Pro  
405 410 415

Ala Lys Leu Met Ala Ala His Ile Thr Met Lys Thr Asn Pro Phe Val  
420 425 430

Trp Ser Ser Cys Ser Arg Asp Tyr Ile Thr Ser Phe Leu Asp Ser Gly  
435 440 445

Leu Gly Leu Cys Leu Asn Asn Arg Pro Pro Arg Gln Asp Phe Val Tyr  
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130 135 140 145 150 155 160 165 170 175 180 185 190 195 200 205 210 215 220 225 230 235 240 245 250 255 260 265 270 275 280 285 290 295 300 305 310 315 320 325 330 335 340 345 350 355 360 365 370 375 380 385 390 395 400 405 410 415 420 425 430 435 440 445 450 455 460

Pro Thr Val Ala Pro Gly Gln Ala Tyr Asp Ala Asp Glu Gln Cys Arg  
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Phe Gln His Gly Val Lys Ser Arg Gln Cys Lys Tyr Gly Val Glu Arg  
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Val Phe Leu Leu Ser Phe Leu Gly Gly Glu Gly Ala Ala Gly Ala Gly  
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Leu Arg Trp Ala Leu Ala Leu Gly Leu Gly Leu Met Phe Glu Val Thr  
 35 40 45

His Ala Phe Arg Ser Gln Asp Glu Phe Leu Ser Ser Leu Glu Ser Tyr  
 50 55 60

Glu Ile Ala Phe Pro Thr Arg Val Asp His Asn Gly Ala Leu Leu Ala  
 65 70 75 80

Phe Ser Pro Pro Pro Pro Arg Arg Gln Arg Arg Gly Thr Gly Ala Thr  
 85 90 95

Ala Glu Ser Arg Leu Phe Tyr Lys Val Ala Ser Pro Ser Thr His Phe  
 100 105 110

Leu Leu Asn Leu Thr Arg Ser Ser Arg Leu Leu Ala Gly His Val Ser  
 115 120 125

Val Glu Tyr Trp Thr Arg Glu Gly Leu Ala Trp Gln Arg Ala Ala Arg  
 130 135 140

Pro His Cys Leu Tyr Ala Gly His Leu Gln Gly Gln Ala Ser Thr Ser  
 145 150 155 160

His Val Ala Ile Ser Thr Cys Gly Gly Leu His Gly Leu Ile Val Ala  
 165 170 175

Asp Glu Glu Glu Tyr Leu Ile Glu Pro Leu His Gly Gly Pro Lys Gly  
 180 185 190

Ser Arg Ser Pro Glu Glu Ser Gly Pro His Val Val Tyr Lys Arg Ser

205

Arg Gln Cys Lys Tyr Gly Val Glu Arg Val Phe Leu Leu Ser Phe Leu  
515 520 525

Gly Gly Glu Gly Ala Ala Gly Ala Gly Gly Gly Asn Ser Thr Arg Ala  
530 535 540

Pro Thr Trp Glu Val Arg Gly Trp Glu Asn Arg Glu Gly Gly  
545 550 555